(1) GENERAL INFORMATION:

- (ii) TITLE OF INVENTION: MODULATION OF HEDGEHOG-MEDIATED SIGNALING PATHWAY
- (iii) NUMBER OF SEQUENCES: 8
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Gray Cary Ware & Freidenrich LLP
 - (B) STREET: 4365 Executive Drive, Suite 1600
 - (C) CITY: San Diego
 - (D) STATE: CA
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 92121
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: Windows95
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE: Herewith
 - (C) CLASSIFICATION: Application
- (vii) PRIORITY APPLICATION DATA:
 - (A) APPLICATION NUMBER: 09/023,249
 - (B) FILING DATE: 2/13/98
 - (C) CLASSIFICATION: Application
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Haile, Lisa A.
 - (B) REGISTRATION NUMBER: 38,347
 - (C) REFERENCE/DOCKET NUMBER: JHU1510-1
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 858/677-1456
 - (B) TELEFAX: 858/677-1465
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18 BASE PAIRS
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: both
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: genomic DNA
 - (ix) FEATURE:
 - (A) NAME/KEY:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 22 BASE PAIRS

(B) LOCATION:	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GTTCTACATA ATGCGCCG	18
(2) INFORMATION FOR SEQ ID NO:2:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 BASE PAIRS (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: genomic DNA	
(ix) FEATURE: (A) NAME/KEY: (B) LOCATION:	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
CGGCGCATTA TGTAGAAC	18
(2) INFORMATION FOR SEQ ID NO:3:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 BASE PAIRS (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GTTCTACGTG ATGCGCCG	18
(2) INFORMATION FOR SEQ ID NO:4:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
TCCCGAAGAC CACCCACAAT GA	22
(2) INFORMATION FOR SEQ ID NO:5:	

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(B) TYPE: nucleic acid(C) STRANDEDNESS: both(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	22
TCAAACCTAC TTTGGGGTCT TA	22
(2) INFORMATION FOR SEQ ID NO:6:	·
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 BASE PAIRS (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	24
ATCGCCGGGG ATGAGCTGGC GGCT	24
(2) INFORMATION FOR SEQ ID NO:7:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 BASE PAIRS (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	27
GATATGGCAA TGGTAGTCAG CACGTGG	2 /
(2) INFORMATION FOR SEQ ID NO:8:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 BASE PAIRS (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	2:
AGCTTCTCCA CTTGCTCTTG G	2.